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                                                                                                                                                                                                                                                                                                                                                The present sequence encodes a protein designated 7F4. This protein is capable of inducing differentiation in osteoblast cells. The protein may be used to screen compounds for the ability to bind to it, for use as ligands, agonists or antagonists and inhibiting or otherwise altering its differentiation inducing activity. Compounds so identified, as well as the protein itself, DNA encoding it, and antibodies to it, may be used in the treatment of diseases of bone growth and osteoblast differentiation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Receptor protein inducing differentiation extracellular region only and can be used
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                                   TTCTGCAGGTACATTTGTCAAGGCGCCCTGCGAAATCCCCCCATACTCAAGGACAATGTGA
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 TCCCTCAGGTACATTTGTCAAGGCGCCCTGCAAAATCCCCCATACTCAAGGACAATGTGA
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                                                                                                                                             Sequence 1509 BP; 387 A; 403 C; 294 G; 425 T; 0 U; 0 Other;
                                                                                                                                                                          The present invention relates to a transgenic non-human animal having the expression of 7F4 gene artificially modified. The transgenic animals are a disease model for bone and glycolipid metabolism disorders. Substances identified by the screening method are agents for the prevention and treatment of diseases including osteoporosis, obesity and diabetes. The present sequence represents the modified mouse 7F4 encoding sequence.
                                                                                                                                                                                                                                                                                                                           Claim
                                                                                                                                                                                                                                                                                                                                            Transgenic non-human animal with modified expression of 7F4 gene screening remedies for bone or glycolipid metabolism disorders.
                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-340227/31.
P-PSDB; ADM46623.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kake T,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7F4 gene; Osteopathic; Anorectic; Antidiabetic;
glycolipid metabolism disorder; osteoporosis; obesity; diabetes; ds.
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                        GCTGCTGCTGCTGCTGCTGAATCTGCCCTTGCAGGTAAAATTTGCTATGCTAGAATT 140
 GTTCCTCTTGCTGCTGCTGCTGAATCTGTTCTTGCCGGTAATATTTGCTATGCTGAATC 109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Saito H,
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                                                                            Score 1182.4; DB 12;
Pred. No. 1.1e-279;
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                                                                                         AGCATTCAAACACATGAGTCTATGAGGGCCAAAACCAATTCAAACCACCACAAGGTTAACAA
                                                                                                                                                             AGCATTCAAACACATGAGTCTATGAGGGCCAAACCAATTCAAACCACCACAGGTTAACAA 1099
                                                                                                                                                                                                              CAGTGACAAACTTCCTCCAACAAGGTCATACCTCCTAATAATACCATTTCTTATGAGGCA 1039
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Novel tmst2-receptor polypeptide useful for diagnosing and treating disease e.g., autoimmune disease, cachexia, cancer or viral, bacterial
                                                        WPI; 2004-224390/21.
N-PSDB; ADJ45749.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
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                                                                                                 Saris C;
                                                                                                                                                 09-JUL-1999;
07-JUL-2000;
                                                                                                                                                                                  17-JUL-2003; 2003US-00622407.
                                                                                                                                                                                                                                                                                        acquired immunodeficiency syndrome; AIDS; anaemia; autoimmune disease; cachexia; cancer; cerebral malaria; diabetes mellitus; disseminated intravascular coagulation; haemorrhagic shock; hepatitis; insulin resistance; lgprosy; leukaemia; lymphoma; meningitis; multiple sclerosis; ischaemia; obesity; organ rejection; rheumatoid arthritis; septic shock; stroke; adult respiratory distress syndrome; ARDS; tuberculosis; viral disease
                                                                                                                          (SARI/)
                                                                                                                                                                                                                  29-JAN-2004.
                                                                                                                                                                                                                                          US2004018544-A1.
                                                                                                                                                                                                                                                                    Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                           Mouse; tmst2-receptor; tmst2 00004-d1;
                                                                                                                                                                                                                                                                                                                                                                                                                   Murine tmst2 00004-d1 polypeptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADJ45750;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADJ45750 standard; protein; 133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and their secreted splice variants, belonging to the tumour necrosis factor (TNP) receptor super gene family and polymucleotides encoding such or ameliorating diseases associated with or resulting from abnormal expression of its putative ligand, such as sepsis, and parasitic diseases, inflammatory diseases, viral, bacterial identification or mapping. The invention is useful in gene therapy. The present sequence is mouse tmst2-receptor protein used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to transmembrane decoy-receptor (tmst2) proteins and their secreted splice variants, belonging to the tunour necrosis
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                                                                                                                        SARIS C.
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                                                                                                                                             99US-0143063P,
2000US-00612033,
                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Š
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      68.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 748; DB 7; ]
; Pred. No. 5.9e-54;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 133,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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                                                                                                                                                                                                                                                                                          disease;
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ID AAWE
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 133; Conserv
            treatment of bone growth
                  Receptor protein inducing differentiation extracellular region only and can be used
                                                        WPI; 1998-568275/48.
N-PSDB; AAV68046.
                                                                                         Kimura N,
                                                                                                           (CHUG-) CHUGAI RES INST
                                                                                                                                   01-APR-1997;
                                                                                                                                                        01-APR-1998;
                                                                                                                                                                               08-OCT-1998.
                                                                                                                                                                                                    WO9843998-A1
                                                                                                                                                                                                                                    Protein
                                                                                                                                                                                                                                                                                         Unidentified.
                                                                                                                                                                                                                                                        Peptide
                                                                                                                                                                                                                                                                                                             Protein 7F4;
                                                                                                                                                                                                                                                                                                                                 Amino acid sequence
                                                                                                                                                                                                                                                                                                                                                         28-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                              AAW80254;
                                                                                                                                                                                                                                                                                                                                                                                             AAW80254 standard; protein; 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a tmst2-receptor polypeptides and the polynucleotide encoding them. The sequences of the invention are useful for treating diseases and conditions including acquired immunodeficiency syndrome (AIDS), anaemia, autoimmune diseases, cachexia, cancer, cerebral haemorrhagic shock, hepatitis, insulin resistance, leprosy, leukaemia, lymphoma, meningitis, multiple sclerosis, ischaemia, obssity, organ distress syndrome (ARDS), tuberculosis and a number of viral diseases. This sequence represents a murine tmst2-receptor polypeptide clone of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 133 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; SEQ ID NO 6; 57pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                          121
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                                                                                        Toyoshima
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   KNCSAGTFVKAPCEIPHTQGQCEKCHPGTFTEKDNYLDACILCSTCDKDQEMVADCSATS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MPGPPCSIVSSISRWPLWRRLLLLLLLLLLLLLVLPLQVKPAWLBLHSPKCPAGEYWSKOVCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MFGFFCSLVSSLSRWFLWRRLLLLLLLLLLLLLNLPLQVKFAMLELHSFKCFAGEYWSKDVCC
                                                                                                                                                                                                                                                                                              differentiation; osteoblast cell; bone growth; bone sarcoma.
                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                              97JP-00099653.
                                                                                                                                                     98WO-JP001511.
                                                                                                                                                                                                                  /note= "mature protein"
                                                                                                                                                                                                                                     /note=_"signal peptide"
                                                                                                                                                                                                                                                             Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                  o
f
                                                                                                     MOLECULAR MEDICINE INC
                                                                                                                                                                                                                                                                                                                               protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           68.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          133
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 748; DB 8; 1
Pred. No. 5.9e-54;
                                                                                                                                                                                                                                                                                                                                7F4.
                                                                                                                                                                                                                                                                                                                                                                                              A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
              in osteoblast cells -
for screening substance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 133;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                substances
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
                for
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           120
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Claim 1; Page 29-31; Slpp; Japanese

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RESULT 10
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Matches 142;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
         Novel ymkz5-receptor polypeptide useful for treating diseases such as tumor, cancer, AIDS, anemia, autoimmune diseases, cachexia, leprosy, leukemia, hepatitis, multiple sclerosis.
                                                                    WPI; 2004-008943/01.
N-PSDB; ADF57550.
                                                                                                                                                    09-JUL-1999; 99US-0143137P
07-JUL-2000; 2000US-00611989
                                                                                                        Zhang K;
                                                                                                                                                                                         11-JUL-2002; 2002US-00193616
                                                                                                                                                                                                                                                                                                          Transmembrane decoy receptor; ymkz5; tumour necrosis factor; cancer; acquired immune deficiency syndrome; AIDS; anaemia; autoimmune disease; cachexia; leprosy; leukaemia; hepatitis;
                                                                                                                                 (ZHAN/)
                                                                                                                                                                                                                                            US2003096355-A1
                                                                                                                                                                                                                                                                    Mus musculus
                                                                                                                                                                                                                                                                                                    multiple sclerosis; myocardial
                                                                                                                                                                                                                                                                                                                                                                       Mouse ymkz5
                                                                                                                                                                                                                                                                                                                                                                                                                        ADF57551;
                                                                                                                                                                                                                                                                                                                                                                                                                                         ADF57551 standard; protein; 176 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence represents a protein designated 7P4. This protein is capable of inducing differentiation in osteoblast cells. The protein may be used to screen compounds for the ability to bind to it, for use as ligands, agonists or antagonists and inhibiting or otherwise altering its the protein inducing activity. Compounds so identified, as well as the protein itself, DNA encoding it, and antibodies to it, may be used in treatment of diseases of bone growth and osteoblast differentiation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 176 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             171
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                                                                                                                                ZHANG K.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TPVKAPCKI PHTQGQCEKCHPGTFTGKDNGLHDCELCSTCDKDQNMVADCSATSDRKCEC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TFVKAPCEIPHTQGQCEKCHPGTFTEKDNYLDACILCSTCDKDQEMVADCSATSDRKCQC 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SIVSSISRWPIWRRLILLILLILINIPLOVKPAMLEIHSPKCPAGEYWSKDVCCKNCSAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SHVSSLSHWF-----LLLLLLLNLFLFVTFAMPESYSFNCPDGEYQSNDVCCKTCPSG
                                                                                                                                                                                                                                                                                                                                                                   receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    68.7%;
75.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 746.5; DB
Pred. No. 1e-53;
8; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8,
                                                                                                                                                                                                                                                                                            ischaemia; obesity; gene therapy; mouse;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
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                                                                                                                                                                                                                                                                                                                                   TNF;
                                                                                                                                                                                                                                                                                                                                   tumour;
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Claim 13;

SEQ ID NO

8

57pp; English

The present invention relates to a transgenic non-human animal having the expression of 7F4 gene artificially modified. The transgenic animals are a disease model for bone and glycolipid metabolism disorders. Substances identified by the screening method are agents for the prevention and

Claim 3; SEQ ID NO 2; 44pp;

Japanese.

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RESULT 11
ADM46623
ID ADM46
XX ADM46
XX ADM46
XX MOUSE
XX MOUSE
XX PF4 9
XX 919co
OS MUS m
XX WO200
XX WO200
XX II-SE
XX II-SE
XX CHUS
XX WPI;
DR SCree
XX Claim
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches 142;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
                                                                                                                                                            Transgenic non-human animal with modified expression screening remedies for bone or glycolipid metabolism
                                                                                                                                                                                                                                                                                                                                 Kake T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-SEP-2003; 2003WO-JP011545.
                                                                                                                                                                                                                                                                                                                                                                                                                            17-SEP-2002; 2002JP-00270321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus
                                                                                                                                                                                                                                                                                                                                                                            (CHUS ) CHUGAI SEIYAKU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2004026026-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7F4 gene; Osteopathic; Anorectic; Antidiabetic; glycolipid metabolism disorder; osteoporosis; o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mouse 7F4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADM46623;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADM46623 standard; protein; 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to transmembrane decoy receptor, ymkz5 belonging to tumour necrosis factor (TNP) receptor supergene family and nucleic acid sequences encoding such receptors. The invention is useful for detecting ymkz5-receptor gene such as tumours or cancers. The sequences of mutated invention are used as medication for a number of diseases such as acquired immune deficiency syndrome (AIDS), anaemia, autoimmune diseases, cachexia, leprosy, leukaemia, hepatitis, multiple sclerosis, myocardial ischaemia, obesity etc. The invention is also useful in gene therapy. The present sequence is mouse ymkz5 receptor protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 176 AA;
                                                                                                                                                                                                                                                  2004-340227/31.
DB; ADM46622.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  171 ---LIVECI 176
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                                                                                                                                                                                                                                                                                                                            Saito H,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7 SIVSSISRWELWRRLILLILLILLILLILARDOVKEAMLELHSEKCEAGEYWSKDVCCKNCSAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Simi)larity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TPVKAPCEIPHTQGQCEKCHPGTFTEKDNYLDACILCSTCDKDQEMVADCSATSDRKCQC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TPVKAPCKIPHTQGQCEKCHPGTFTGKDNGLHDCELCSTCDKDQNMVADCSATSDRKCEC
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                                                                                                                                                                                                                                                                                                                         Makishima
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    68.7%;
75.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 746.5;
Pred. No. 1e-5
8; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      osteoporosis; obesity; diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1e-53;
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                                                                                                                                                          of 7F4 gene
disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence encodes a protein designated 7F4. This protein is capable of inducing differentiation in osteoblast cells. The protein may be used to screen compounds for the ability to bind to it, for use as ligands, agonists or antagonists and inhibiting or otherwise altering its differentiation inducing activity. Compounds so identified, as well as the protein itself, DNA encoding it, and antibodies to it, may be used in the treatment of diseases of bone growth and osteoblast differentiation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Receptor protein inducing differentiation extracellular region only and can be used treatment of bone growth disorders.
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                                                                                    ATGCGAGTGCCAAATAGGTCTTTACTACTATGACCCAAAATTTTCCGGAATCATGCCGCCC
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                                            GAAGTGTCACCCAGGAACATTCACAGGGAAAGATAATGGCCTGCATGATTGTGAACTTTG
                                                                                                                                                            TTCTGCAGGTACATTTGTCAAGGCGCCCTGCGAAATCCCCCCATACTCAAGGACAATGTGA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-JUN-2004
                                                                                                                                                                                                                                                                                                                                                              Sequence 176 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 3; SEQ ID NO 2; 44pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CHUS ) CHUGAI SEIYAKU KK.
                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2004-340227/31.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TFVKAPCEIPHTQGQCEKCHPGTFTEKDNYLDACILCSTCDKDQEMVADCSATSDRKCQC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SHVSSLSHWF----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SLVSSLSRWFLWRRLLLLLLLLLLLLPLQVKFAMLELHSFKCPAGEYWSKDVCCKNCSAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           QIGLYYYDPKFPESCRPCTKCPQGIPVLQECNSTANTVCSSSVS
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                                                                             TFVKAPCEIPHTOGOCEKCHPGTFTEKDNYLDACILCSTCDKDQEMVADC$ATSDRKCQC
  RTGLYYYDPKFPESCRPCTKCPQGIPVLQECNSTANTVCSSSVS 170
                                                     TFVKAPCKIPHTQGQCEKCHPGTPTGKDNGLHDCELCSTCDKDQNMVADCSATSDRKCEC
                                                                                                                                                                                                     SIVSSISRMFLWRRLLLLLLLLLLLLLPLQVKFAMLELHSFKCPAGEXWSKDVCCKNCSAG
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                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                   69.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 79.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          disorder; osteoporosis; obesity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Anorectic; Antidiabetic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -LLLLLLLILFLPVIFAMPESYSFNCPDGEYQSNDVCCKTCPSG
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5; Mismatches 20;
                                                                                                                                                          LLLLLLNLFLPVIFAMPESYSFNCPDGEYQSNDVCCKTCPSG
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                                                                                                                                                                                                                                                           Score 705; DB 8; ]
Pred. No. 2.1e-50;
5; Mismatches 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   expression of 7F4 gene metabolism disorders.
                                                                                                                                                                                                                                                                                                             Length 176;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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RESULT 13
ADM46624
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                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                   The present invention relates to a transgenic non-human animal having the expression of 7F4 gene artificially modified. The transgenic animals are a disease model for bone and glycollpid metabolism disorders. Substances identified by the screening method are agents for the prevention and treatment of diseases including osteoporosis, obesity and diabetes. The present sequence represents the modified mouse 7F4 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO2004026026-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7F4 gene; Osteopathic; Anorectic; Antidiabetic; glycolipid metabolism disorder; osteoporosis; obesity; diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mouse 7F4 protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-JUN-2004
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                                                                                                                                                                                                                                                                                                                                Sequence 148 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 3; SEQ ID NO 3; 44pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tranagenic non-human animal with modified screening remedies for bone or glycolipid
                                                 ABG09344;
                                                                       ABG09344
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                                                                                                                                                                                                       ACILCSTCDKDQEMVADCSATSDRKCQCRTGLYXYDPKFPESCRPCTKCPQGIPVLQECN
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                                                                                                                                                                                DCBLCSTCDKDQNWVADCSATSDRKCBCQIGLYYYDPKFPBSCRPCTKCPQGIPVLQBCN
                                                                                                                                                                                                                                AMPESYSFNCPDGEYQSNDVCCKTCPSGTFVKAPCKIPHTQGQCEKCHPGTFTGKDNGLH
                                                                                                                                                                                                                                               AMLELHSFKCPAGEYMSKDVCCKNCSAGTFVKAPCEIPHTQGQCEKCHPGTFTEKDNYLD
                                                                         standard;
                                                                                                                                                        STANTYCSSSVS 170
                                                                                                                                   STANTVCSSSVS 132
                                                                                                                                                                                                                                                                                 Conservative
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                                                                       protein;
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Pred. No. 3.3e-45;
5; Mismatches 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    expression of 7F4 gene metabolism disorders.
                                                                                                                                                                                                                                                                                                          Length 148
                                                                                                                                                                                                                                                                                    Indels
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Novel human diagnostic protein #9335.

13-FEB-2002

(first entry)

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